

REPLACEMENT SHEET

16/61

Sequences of the (+) strand

complete forms:

Seq.	ID
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	B11
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	7
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	AAAACU---UAAGUCCGAAGGUAACCAUCUAC--
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	ACGUACCACUGUCGGGUCCAC
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	G2
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	AAAACU---UAAGUCCGAAGGUAACCAUCUAC--
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	ACGUACCACUGUCGGGUCCAC
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	E12
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	AAAACU---UAAGUCCGAAGGUAACCAUCUAC--
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	ACGUACCACUGUCGGGUCCAC
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	B7
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	AAAACG---UAAGUCCGAAGGUAACCAUCUAC--
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	ACGUACCACUGUCGGGUCCAC
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	A8
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	AAAACU---UAAGUCGUAGGUAACCAUCUAC--
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	ACGUACCACUGUCGGGUCCAC
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	B12
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	AAAACU---UAAGUCGUAGGUAACCAUCUAC--
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	ACGUACCACUGUCGGGUCCAC
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	E3
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	AAAAC---GUAAAUCGGAGGUAACCAUCUAC--
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	ACGUACCACUGUCGGGUCCAC
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	C12
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	AAAAC---GUAAAUCGGAGGUAACCAUCUAC--
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	ACGUACCACUGUCGGGUCCAC
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	C11
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	AAAAC---GUAAAUCGGAGGUAACCAUCUAC--
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	ACGUACCACUGUCGGGUCCAC
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	A3
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	AAAAC---GUAAAUCGGAGGUAACCAUCUAC--
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	ACGUACCACUGUCGGGUCCAC
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	F5
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	AAAAC---GUAAAUCGGAGGUAACCAUCUAC--
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	ACGUACCACUGUCGGGUCCAC
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	A12
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	29
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	AAAAC---GUAAAUCGGAGGUAACCAUCUAC--
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	ACGUACCACUGUCGGGUCCAC
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	F12
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	31
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	AAAAC---GUAAAUCGGAGGUAACCAUCUAC--
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	ACGUACCACUGUCGGGUCCAC
GGAGCUCAGACUUACUCUGUGG---AGGCCAUU	G5

GGAGCUCAGACUUACUCUG
DE. 40F-Primer (Seq. ID No. 35)

Primer moieties underlined and in bold

CGUACCACUGUCGGGUCCAC
DE. 40R-Primer (rev. und
compl.) (Seq. ID No. 36)

Fig. 13-1